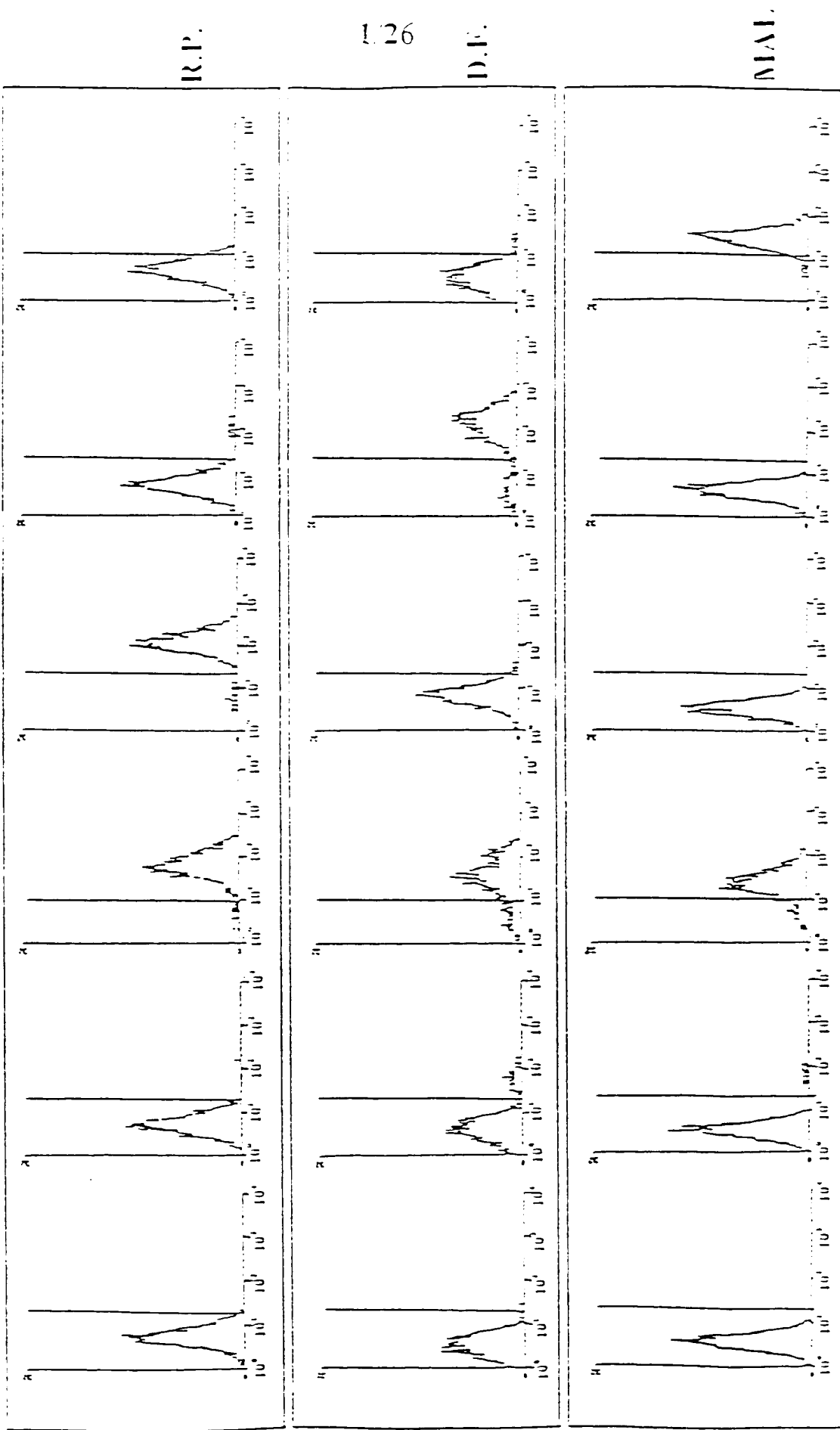


C anti-CD3 anti-CD16 anti-CD158 anti-CD158 anti-CD158
 (p50.1, EB6) (p50.2, GL183) (p50.3, PAX250)

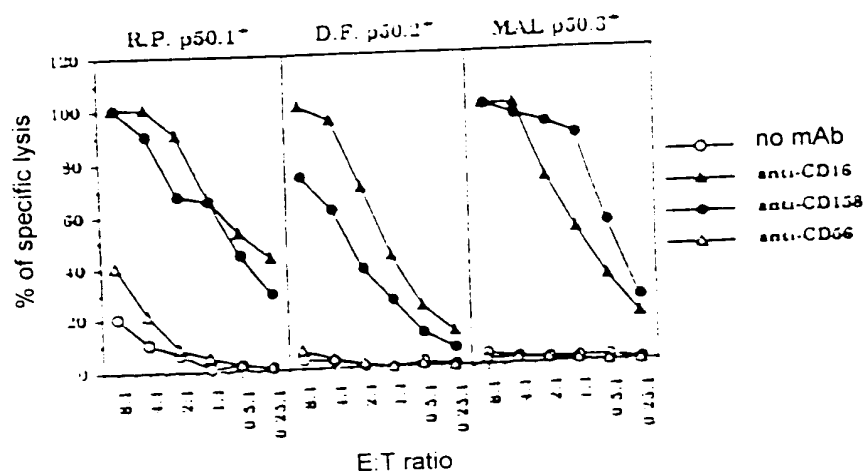


Intensity of fluorescence

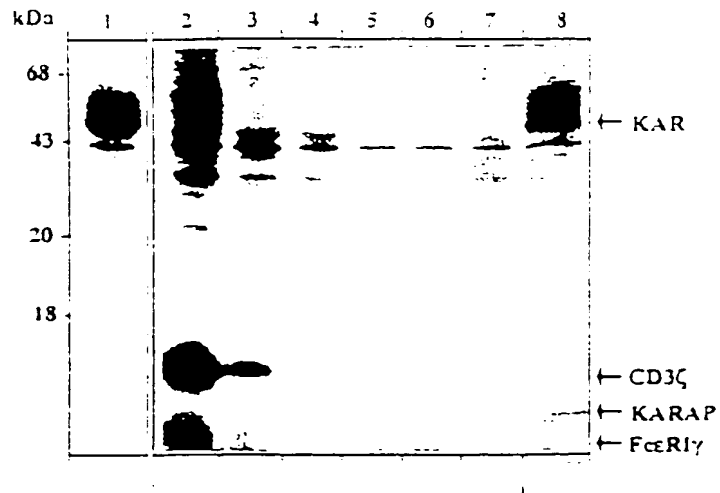
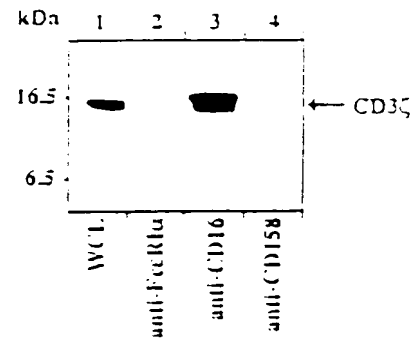
Figure 1A

2.26

Figure 1B



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Figure 2AFigure 2B

426

Figure 3A

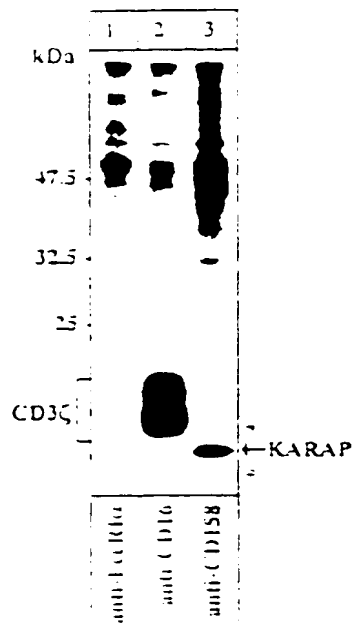


Figure 3B

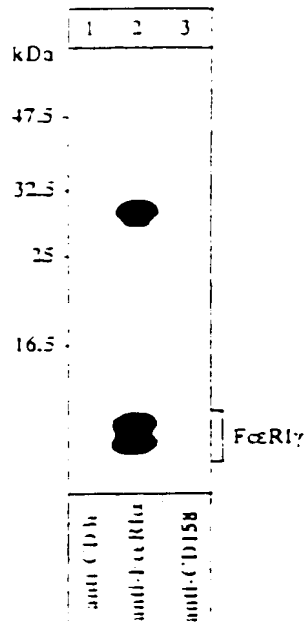
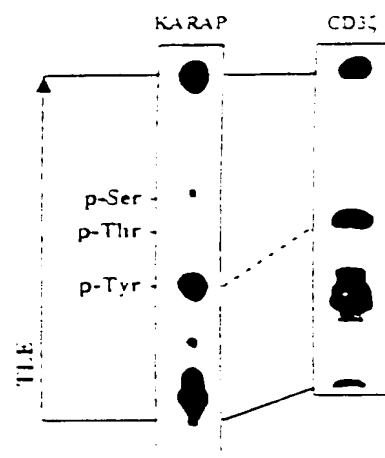
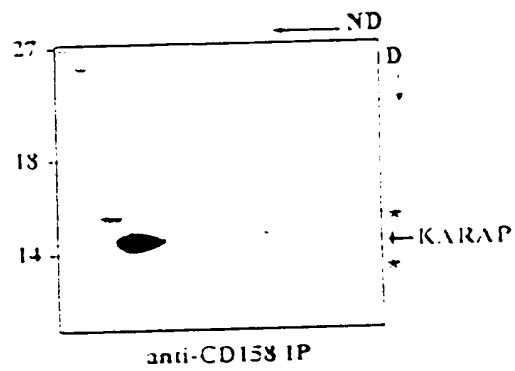


Figure 3C

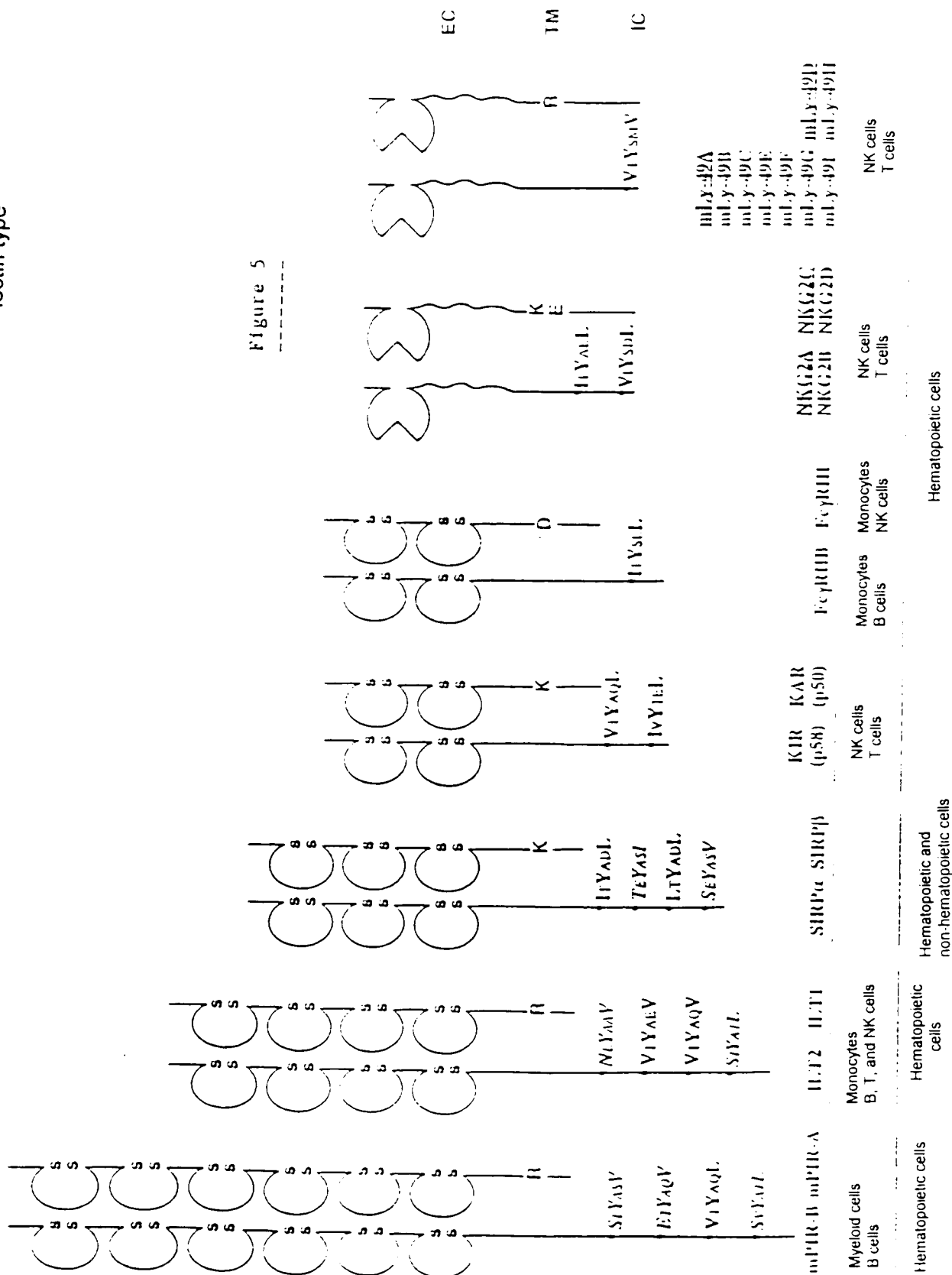


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Figure 4

lectin type

Figure 5



NK p58/50 cell receptors for class I MHC molecules

Figure 6

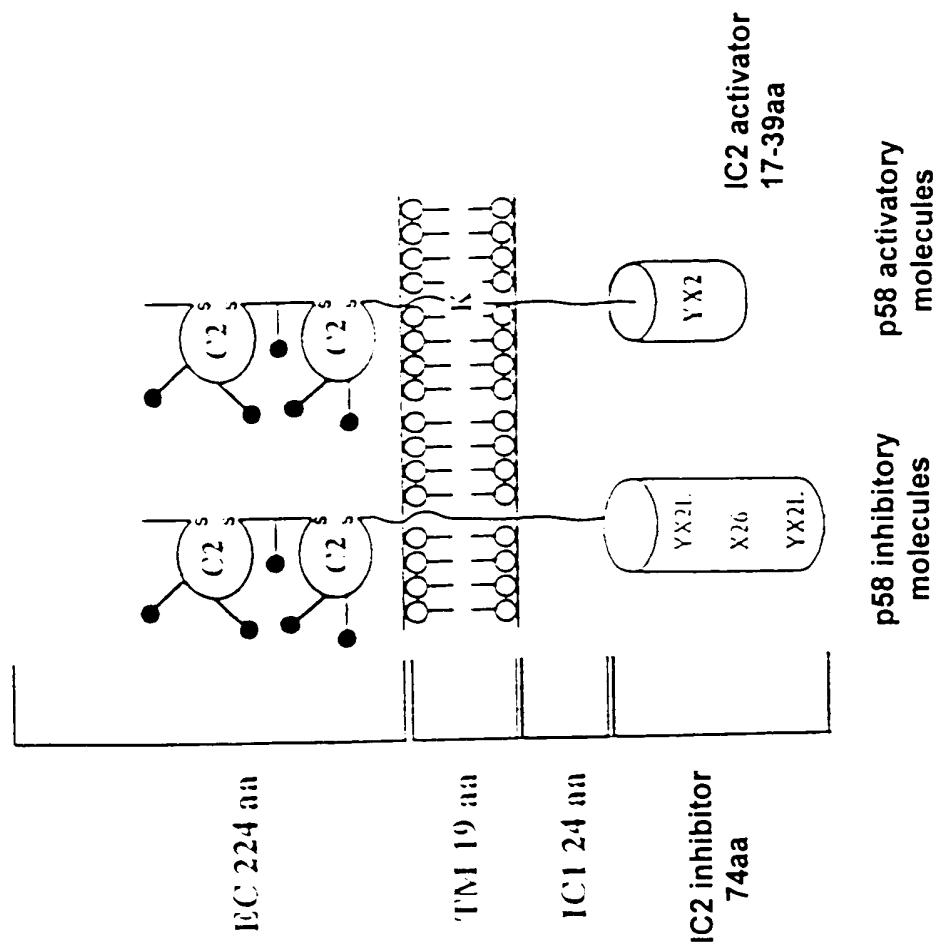


Figure 7

1 ggtaacacaa ggtccacaa gccctggac tgggtgtac agtgcatac tggccacaa:
61 ggggtctgg agctctgg tgcctgt tctctgt cctctgact gtgggaggat
121 taagtccgt acaggccag agtgacac tccaagatg cgaatgtct tccgtgagcc
181 ctgggtact gtctgggaa gttctgggt acttggtg gactctgtg attgccctg
241 ctgtgtact tctggccgc ctggtctcc gaggtaagg gacagcggaa gggaccggg
301 aacaacacat tctgagact ggtctgct atcaggagct tcagggtcag agacatgaag
361 tatacgtga cttcaacaa caggggcaat attacagatg agccactct atgccatca
421 ggggtctgat gccaggatcc ggtcaaca gatgctact caacaagccc tctctgagat
481 caggactccc gttgaatac agatccacag ggtacct

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Fluorene 3

1/1

cag agt gac acc ttc cca aga tgc gac agt ttc ttc gtc agc ccc ggt gta ctc ttc ggt
 Q S D T F P R C C C S S V S P G V L S G

31/11

61/21

31/31

att gtt ctc ggt gac ttc gtc ttc acc ctc ctc att gcc ctc gcc gtc ttc ttc ctc ggc
 I V L G D L V L T L L L A L A V Y S L G

121/41

151/51

cgc ctc gtc ttc cga ggt cta ggt acc gtc gaa ggt acc cgc aaa cta ctc att gcc gac
 R L V S R G Q G T A E G T R K Q H I A E

131/61

211/71

att gac ctc ctc ttc ctc gtc ctc ctc ggt ctc gaa ctc gaa gta ttc agt gtc ctc att
 T E S P Y Q E L L G Q R H E V Y S D L N

241/81

acc cag agt cta ttc ttc aga

T Q R Q Y Y R

Figure 3

ITAM polypeptides

CD3 ζ_1	YneLnlgrrre-YcrL
CD3 ζ_2	YneLgkckmaaaYseL
CD3 ζ_3	Yqgls tatkat-YdaL
CD3 γ	Yqplkoreccq-YsnL
CD3 δ	Yqplrcrrccq-YshL
CD3 ϵ	YepLrkqqrnl-YsgL
I $\epsilon\alpha$ (CD79a)	Yedls cglqgt-YqcV
I $\epsilon\beta$ (CD79b)	Yeqldl cqtat-YedL
Fc ϵ R γ	Yqplcttrnges-YetL
Fc ϵ R β	YeeLmlysat--YsaL
Fc ϵ R α	YqaLqggqrneY-YsdL
Consensus	Y--L-----Y--L

Figure 10A

SEQ ID n°6

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
TCACACCGAG	TTCGACGAG	CGTCGACTG	TGGTGTCCAG	TGCATATCTG	50
GGCAGCATCG	GCCTGTGAG	CGTCGTGGG	CGTTGTGTTG	CTTCTGTCTCC	100
TCTGACTCT	GGCAGCTTA	ACTCCGTGAC	ACGCGTGCAG	TGACACTTTC	150
CGACGATTCG	ATTCTTCTTC	CGTCGCGCT	GGTGTACTGT	CTGGGATGT	200
TCTGGGTGAC	TGGTGTGTA	CTCTCTGAT	TGCGCTGGCT	GTGACTCTTC	250
TGGCGCGCT	GGTGTCCGA	GGTCACCGA	CGCGGTGAG	GACCGCGAA	300
CGACGCTTC	CTGCTCTTA	GTGCGTTTC	CGCGGTTTC	AGGCTCTTC	350
ACTGCTCTA	TACCTGCTC	TGACGCTCA	CGCGCTATC	TACGCTCTG	400
CGCTCTCTC	GGCTCTCTC	GGCTCTCTC	CGCGCTCTC	TGCTCTCTC	450
TGCTCTCTC	AGCTCTCTC	TGCTCTCTC	CGCTCTCTC	TGCTCTCTC	500
ATCTCTCTC	CTCTCTCTC				515

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
SHYPPAPE	WCFVHNEPW	GGGAWCLAF	LPVLTUCEL	SPVQACSDTF	50
FPDCCSVSP	GVLSGIVLGD	WVTLILALA	VYSLGELVSR	CGGREGIRK	100
GVPAIESPA	GVLSGQFHEV	YSILNTQFCY	WVGFHEPIS	GVMPGGGSR	150
GVLPFSIRS	GVLEVFSTG	Y			171

Figure 10B

SEQ ID n°11

12.26

Figure 11A

SEQ ID n°7

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
GTGCGCTCTT	GGGCGCGCTG	GGGCGCTCTG	AGCGCTCTCT	GTGCGCTCTG	50
TTGCGCTCTG	TTGCGCTCTG	TTGCGCTCTG	TTGCGCTCTG	TTGCGCTCTG	100
GTGCGCTCTT	TTGCGCTCTG	GGGCGCTCTG	TTGCGCTCTG	GTGCGCTCTG	150
TTGCGCTCTG	TTGCGCTCTG	GGGCGCTCTG	TTGCGCTCTG	GTGCGCTCTG	200
GTGCGCTCTT	GGGCGCTCTG	GGGCGCTCTG	GGGCGCTCTG	GGGCGCTCTG	250
GGGCGCTCTG	GGGCGCTCTG	GGGCGCTCTG	GGGCGCTCTG	GGGCGCTCTG	300
TTGCGCTCTG	GGGCGCTCTG	GGGCGCTCTG	GGGCGCTCTG	GGGCGCTCTG	350
TTGCGCTCTG	GGGCGCTCTG	GGGCGCTCTG	GGGCGCTCTG	GGGCGCTCTG	371

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
ATGCGCTCTT	GGGCGCTCTG	GGGCGCTCTG	GGGCGCTCTG	GGGCGCTCTG	50
ATGCGCTCTT	GGGCGCTCTG	GGGCGCTCTG	GGGCGCTCTG	GGGCGCTCTG	100
ATGCGCTCTT	GGGCGCTCTG	GGGCGCTCTG	GGGCGCTCTG	GGGCGCTCTG	123

Figure 11B

SEQ ID n°12

1326

Figure 12A

SEQ ID n°8

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
GCCTTCGCTT	CCCTTCCTCTC	CTCTCTGACTC	TGGGAGGCTT	AACCTCCCGTA	50
CGAGCCCGCA	GTGAGCTTTT	CCGACAGTCC	CCCTCTTCTT	CCGTGAGCCC	100
TGCTTACTG	CGTGGGCTG	TTTGGGCTG	CTTGGGCTG	AATCTGCTGA	150
TTGCTTGGC	TGTGACTCT	CTGGCCCGCC	TGCTTCCCG	AGCTGAGGG	200
AGAGCCGAG	CGAGCCGGA	AGACACATT	CTTGAGCTG	AATCGCTTA	250
TGAGGCTT	CGGGCTGCA	GACTGACT	AGAGCTGAC	CTCAAGCTC	300
AGAGCTGCA	TTAGCTGCA	CCCTCTCTA	TGCTCTGAC	CGGCTGCTG	350
CGGCTGCTG	GTCTTCTGAC	ATCTCT			376

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
ATCTCTCTA	CGCTTCTGAC	CTCTCTCTA	CTCTCTCTA	CTCTCTCTA	50
ATCTCTCTA	CTCTCTCTA	CTCTCTCTA	CTCTCTCTA	CTCTCTCTA	100
ATCTCTCTA	CTCTCTCTA	CTCTCTCTA	CTCTCTCTA	CTCTCTCTA	124

Figure 12B

SEQ ID n°13

14.2b

SEQ ID n°9

Figure 13A

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
CCAGGCGGCTG	GAATGTGCTG	TGCTGTGCTG	AGTGGGCGAC	GTGGGGGGCT	50
GTGGAGGCTG	GTGGTGGCTG	GTGGTGGCTG	GTGGTGGCTG	GTGGTGGCTG	100
GGATTAAGTG	GGATTAAGTG	GGATTAAGTG	GGATTAAGTG	GGATTAAGTG	150
TGTTGGGCTG	AGGGGCTG	AGGGGCTG	AGGGGCTG	AGGGGCTG	200
TGTTGGGCTG	AGGGGCTG	AGGGGCTG	AGGGGCTG	AGGGGCTG	250
TGTTGGGCTG	AGGGGCTG	AGGGGCTG	AGGGGCTG	AGGGGCTG	300
TGTTGGGCTG	AGGGGCTG	AGGGGCTG	AGGGGCTG	AGGGGCTG	350
TGTTGGGCTG	AGGGGCTG	AGGGGCTG	AGGGGCTG	AGGGGCTG	400
TGTTGGGCTG	AGGGGCTG	AGGGGCTG	AGGGGCTG	AGGGGCTG	402

11

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
25LGGGQQCT	SGHGGGGS	WGLFLPALL	TGGLSPVQA	QSDIFFPQDC	50
SSVSPGVLAG	TLGLVMTL	LPLVYSLS	PLVSPQQT	EGVFPQFLAE	100
TSPVQELQG	QFPPVYSLS	TSPVQELQG	LCP		150

Figure 13B

SEQ ID n°14

15.20

Figure 14A

SEQ ID n°10

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
CTTCTCTCTCT	GTCTCTCTCTA	CTCTCTCTCTG	ATTCTCTCTCT	CTTCTCTCTCT	50
AGCTCTCTCT	TTCTCTCTCTA	TCTCTCTCTCT	CTTCTCTCTCT	CCCTCTCTCTA	100
CTCTCTCTCTA	TTCTCTCTCTG	TCTCTCTCTCT	TTCTCTCTCTG	TCTCTCTCTCT	150
CTCTCTCTCTA	TCTCTCTCTCT	CTCTCTCTCT	CTCTCTCTCTA	GGCTCTCTCT	200
AGCTCTCTCT	CTCTCTCTCTA	ATTCTCTCTCT	CTCTCTCTCT	TCTCTCTCTCT	250
CTCTCTCTCT	AGCTCTCTCTA	AGCTCTCTCT	CTCTCTCTCTA	CTCTCTCTCT	300
AGCTCTCTCT	TCTCTCTCTCT	CTCTCTCTCT	CTCTCTCTCT	AGCTCTCTCT	350
CTCTCTCTCT	CTCTCTCTCTA	CTCTCTCTCT	CTCTCTCTCT	CTCTCTCTCT	400
CTCTCTCTCTA	TCTCTCTCTCT	CTCTCTCTCT	CTCTCTCTCT	TCTCTCTCTCT	450
TCTCTCTCTCT	CTCTCTCTCTA	AGCTCTCTCT	CT		460

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
CTCTCTCTCT	CTCTCTCTCT	CTCTCTCTCT	CTCTCTCTCT	CTCTCTCTCT	50
CTCTCTCTCT	CTCTCTCTCT	CTCTCTCTCT	CTCTCTCTCT	CTCTCTCTCT	100
CTCTCTCTCT	CTCTCTCTCT	CTCTCTCTCT	CTCTCTCTCT	CTCTCTCTCT	150
CTCTCTCTCT	CTCTCTCTCT	CTCTCTCTCT	CTCTCTCTCT	CTCTCTCTCT	160

Figure 14B

SEQ ID n°15






Figure 15 (contd.)

AA098506	-----	-----	-----	-----	-----	402
AA242315	-----	-----	-----	-----	-----	448
W88159	-----	-----	-----	-----	-----	369
AA734769	-----	-----	-----	-----	-----	371
W41142	-----	-----	-----	-----	-----	362
Consensus	AGCCGACTGT	ATGCCCATCA	CCGCCCTGAT	CCCGGCAATC	GGTCAATGCA	450
AA098506	-----	-----	-----	-----	-----	402
AA242315	-----	-----	-----	-----	-----	497
W88159	-----	-----	-----	-----	-----	376
AA734769	-----	-----	-----	-----	-----	371
W41142	-----	-----	-----	-----	-----	412
Consensus	GAAGGCTACT	CAACAAGCCC	TTCTSTGRGA	TCAGGACTCC	CGTGGGACAA	500
AA098506	-----	-----	-----	-----	-----	402
AA242315	-----	-----	-----	-----	-----	515
W88159	-----	-----	-----	-----	-----	376
AA734769	-----	-----	-----	-----	-----	371
W41142	-----	-----	-----	-----	-----	462
Consensus	CAGATCCACA	GGTACCTCC	CTGAGATATC	TCAGATTGTA	CCATTTCCTT	550
AA098506	-----	-----	-----	-----	-----	402
AA242315	-----	-----	-----	-----	-----	515
W88159	-----	-----	-----	-----	-----	376
AA734769	-----	-----	-----	-----	-----	371
W41142	-----	-----	-----	-----	-----	482
Consensus	CCCCAATAG	AAGACCGACA	-----	-----	-----	570

11-25-10

SEQ ID n°11	AA0442015	protein
SEQ ID n°13	W68169	protein
SEQ ID n°15	W41142	protein
SEQ ID n°14	AA098505	protein
SEQ ID n°12	AA034769	protein
SEQ ID n°17	Consensus	

2214211 protein
 2214211 protein
 2214211 protein
 2214211 protein
 2214211 protein
 consensus

 3A4271 protein
 3B3139 protein
 3A2140 protein
 3A3935 protein
 3B3459 protein
Consensus

32042015 protein
 320139 protein
 320142 protein
 3203805 protein
 3204789 protein
 320513

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SRQVFPAPOL  WCPVHINPPA  OSCAS .....
.....
.....
.....
QPLDCCVQCE  SCIRHC .....  OSCAS .....
XPL .....  XPLHC .....  ALPP .....
.....  .....  .....  WCPVHINPPA  SCIRHC  SCIRHC

```

PROCESSING STATION NUMBER 1234567890

[illegible][illegible]

1928

Page 17

SEP 10 1977

10	20	30	40	50	
1014567890	1014567890	1014567890	1014567890	1014567890	
ADACACGCTC	CGACGACGCG	CTGACGCTGC	GTGACGCTGC	GTATGCTGCG	50
CACTATGCGG	CGTTTGACGC	CGTTTGCTGC	CGTTTGCTGC	CTTCTGCTGC	100
TGCTGACTGT	CGAGGTGACT	TGCGCGGCTT	CTTGCGCTGC	CTTCTGCTGC	150
CTTACGCTGT	CTTGCGGCGA	CGCTAGCGCT	CGAGCGACGA	ACGGATACGA	200
CGAGACGCGG	CGAGGCTGCG	CGAGACGAGG	TTCTGTACGA	CGCTGTGCGT	250
TTTACGCTGT	CGTACGCGCG	CTGACGCTGC	CGAGCTGCGA	GTATGCTGAC	300
CGCGGACGCT	CGAGCTGCGA	CGAGACGCTA	CGAGCTGCTG	GTATGCTGCG	350
CGCGGCTGCG	CGTGTGCTGC	CGAGCTGCTG	CGAGCTGCTG	CGAGCTGCTG	400
ACGCTGCTGT	TGCGCTGCGA	TTTACGCTGT	TGTTTGCTGT	ACGCTGCTGC	450
ACGCTGCGCG	CGCTGCTGCT	AAATGCTGCT	CGCTGCTGAC	CGCTGCTGCT	500
ACGCTGCTGT	TGCTGCTGCT	CGCTGCTGCT	AGCTGCTGCT	TTTCTGCTGT	550
CTTACGCTGT	AGCTGCTGAC	AGCGGCTGCT	TGCTGCTGCT	TTTACGCGCG	600
CGCTGCTGCT	TGCTGCTGCT	AAAGCTGCTG	TGCGGCTGCT	CGCTGCTGCT	650
CGCTGCTGCT	TTTCTGCTGAC	ACGCTGCTGCT	CGCTGCTGAC	TTTCTGCGCG	700
CTTCTGCTGCT	TGCTGCTGCT	CTTCTGCTGAC	AGCGGCTGCT	TTTCTGCTGCT	750
CGCTGCTGCT	AGCTGCTGCG	AGCTGCTGCG	GTGCTGCTGCT	CTTCTGCTGCT	800
CTGCTGCTGCT	TGCTGCTGCT	CTTCTGCTGCG	CGCTGCTGCT	CGCTGCTGCT	850
AGCTGCTGAC	AGCTGCTGCT	AGCTGCTGCT	AAATGCTGCT	CGCTGCTGCT	900
CGCTGCTGCT	TGCTGCTGCT	AGCTGCTGCT	CGCTGCTGCT	CTTCTGCTGCT	950
CGCTGCTGCT	CGCTGCTGCT	AGCTGCTGCT	CGCTGCTGCT	GTAGCTGCTG	1000
CGCTGCTGCT	TGCTGCTGCT	CTTCTGCTGCG	CGCTGCTGCT	TTTCTGCTGCT	1050
CGCTGCTGCT	AGCTGCTGCT	AGCTGCTGCT	AGCTGCTGCT	AGCTGCTGCT	1100
AGCTGCTGCT	TGCTGCTGCT	AGCTGCTGCT	AGCTGCTGCT	AGCTGCTGCT	1150
AGCTGCTGCT	AGCTGCTGCT	AGCTGCTGCT	CGCTGCTGCT	AGCTGCTGCT	1200
AGCTGCTGCT	AGCTGCTGCT	CGCTGCTGCT	AGCTGCTGCT	CGCTGCTGCT	1250
TTTCTGCTGCT	CGCTGCTGCT	AGCTGCTGCT	GTGCTGCTGCT	TTTCTGCTGCT	1300
AGCTGCTGCT	TTTCTGCTGCT	CGCTGCTGCT	AGCTGCTGCT	CGCTGCTGCT	1350
TTTCTGCTGCT	TGCTGCTGCT	TGCTGCTGCT	TGCTGCTGCT	TGCTGCTGCT	1400
TGCTGCTGCT	TGCTGCTGCT	TGCTGCTGCT	TGCTGCTGCT	TGCTGCTGCT	1450
CGCTGCTGCT	CGCTGCTGCT	TGCTGCTGCT	AGCTGCTGCT	TTTCTGCTGCT	1500
CTGCTGCTGCT	TGCTGCTGCT	CGCTGCTGCT	TGCTGCTGCT	GTGCTGCTGCT	1550
AGCTGCTGCT	AGCTGCTGCT	CGCTGCTGCT	CGCTGCTGCT	CTTCTGCTGCT	1600
CTTCTGCTGCT	TGCTGCTGCT	TGCTGCTGCT	AGCTGCTGCT	TGCTGCTGCT	1650
CGCTGCTGCT	TTTCTGCTGCT	CGCTGCTGCT	AGCTGCTGCT	AGCTGCTGCT	1700
CGCTGCTGCT	CGCTGCTGCT	CGCTGCTGCT	CTTCTGCTGCT	AGCTGCTGCT	1750
CGCTGCTGCT	TGCTGCTGCT	TGCTGCTGCT	TGCTGCTGCT	CGCTGCTGCT	1800
AGCTGCTGCT	TGCTGCTGCT	TTTCTGCTGCT	TGCTGCTGCT	TTTCTGCTGCT	1850
CGCTGCTGCT	CGCTGCTGCT	AGCTGCTGCT	CGCTGCTGCT	TTTCTGCTGCT	1900
AGCTGCTGCT	AGCTGCTGCT	TGCTGCTGCT	AGCTGCTGCT	CGCTGCTGCT	1950
TGCTGCTGCT	TTTCTGCTGCT	AGCTGCTGCT	TGCTGCTGCT	AGCTGCTGCT	2000
CTTCTGCTGCT	AGCTGCTGCT	TGCTGCTGCT	TGCTGCTGCT	TGCTGCTGCT	2050

20.2b

Figure 17 (contd.)

SEQ ID n° 18 (contd.)

10	20	30	40	50	
1034567890	1034567890	1034567890	1034567890	1034567890	
GGGCTACCA	AACTAACCT	CTCTACAGC	AACTAACCA	ACAAAAACA	2100
AACTAACCA	AACTAACCT	CACTACAGC	CTCTACAGT	CGCTACAGC	2150
CGGCTACCA	TAATTAACA	CACTACAGC	AACTAACCA	AACTAACCA	2200
CTCTACAGT	TAATTAACA	TAATTAACA	GGGCTACCA	TGGCTACCA	2250
CACTACAGC	AACTAACCA	CTCTACAGC	CGCTACAGC	AACTAACCA	2300
GGGCTACCA	TAATTAACA	TAATTAACA	CGCTACAGC	GGGCTACCA	2350
CACTACAGC	AACTAACCA	AACTAACCA	AACTAACCA	CTCTACAGC	2400
GGGCTACCA	AACTAACCA	GGGCTACCA	TGGCTACCA	GGGCTACCA	2450
CTCTACAGT	TAATTAACA	GGGCTACCA	CTCTACAGC	AACTAACCA	2500
GGGCTACCA	TGGCTACCA	AACTAACCA	AACTAACCA	TTTCTACCA	2550
TTTCTACCA	AACTAACCA	CACTACAGC	CACTACAGC	AACTAACCA	2600
AACTAACCA	AACTAACCA	GGGCTACCA	TGGCTACCA	TTTCTACCA	2650
TTTCTACCA	TGGCTACCA	CACTACAGC	CACTACAGC	AACTAACCA	2700
AACTAACCA	TAATTAACA	CACTACAGC	TGGCTACCA	AACTAACCA	2750
TGGCTACCA	GGGCTACCA	AACTAACCA	TGGCTACCA	CTCTACAGC	2800
TGGCTACCA	GGGCTACCA	CACTACAGC	GGGCTACCA		2850

21.26

Figure 133' Intron sequence
(donor site)

Exon sequence

5' Intron sequence
(acceptor site)MetGly aIGluG
ATGCGG...-Exon 1-...TCCAGG

GAGGTGA....

....TCCTTAG

IyLeuS InSerA
GATTAA...-Exon 2-...ACAGTG

GTAAGCC....

....TCCTCAG

spThrP InGluA
ACACTT...-Exon 3-...AACAGA

GTAAGAA....

....TCTCTAG

IyThrA TyrGln
GGACCC...-Exon 4-...TATCAG

GTAAGAA....

....TTTAAAG

Figure 2

22.26

	1034557927	1034557928	1034557929	1034557930	1034557931	
SEQ ID n°27	ATGGGGGGCTT	TGGAGCCCTC	CTGGTGGCTT	CTCTTGGCTT	CTGTGCTGCT	50
SEQ ID n°28	M G A L E P S	N C L L F L E	V L L			
	GACTGTGGAC	GCATTAAGTC	CCCTACACGC	CCACGTGTAC	ACTTTCCCAA	100
	T V E G L S P	V Q A Q S D	T F P R			
	GATGCGACTC	TTCTTCCCTC	AGCCCTGGTC	TACTTGGCTC	CAATTCTTCT	150
	C D C S S V	S P G V L A G	I V L			
	CGTCACTTGG	TCTTCACTCT	CGTCATTGCC	CTCCCTCTCT	ACTCTCTGGG	200
	G D L V L T L	L I A L A V Y	E L S			
	CGCCCTGGTC	TCCCGAGGTC	AACACAGGAC	CGGCAACAA	CACATTGCTC	250
	R L V S R G Q	E R T R K Q	H I A E			
	AGACTGAGTC	GCCTTATCAG	CAGTTTCAGG	GTCAACAGACA	TGAAGTATAC	300
	T E S P Y Q	E L Q G Q R H	E V Y			
	AGTGACCTCA	ACACACAGAG	GCAATATTAC	AGATGAGCCC	ACTCTATGCC	350
	S D L N T Q R	Q Y Y R L A H	S M P			
	CATCAGCGCC	CTGATGCCCC	GATCCGGTCA	TTCACAGATC	CTACTCAACA	400
	I S G L M P G	S G H S R C	L L N K			
	AGCCCTCTCT	GAGATCAGGA	CTCCCGTTGG	AATACAGATC	CACAGGGTAC	450
	P S L R S G	L P L E Y R S	T G Y			
CT						452

23.26

Figure 20

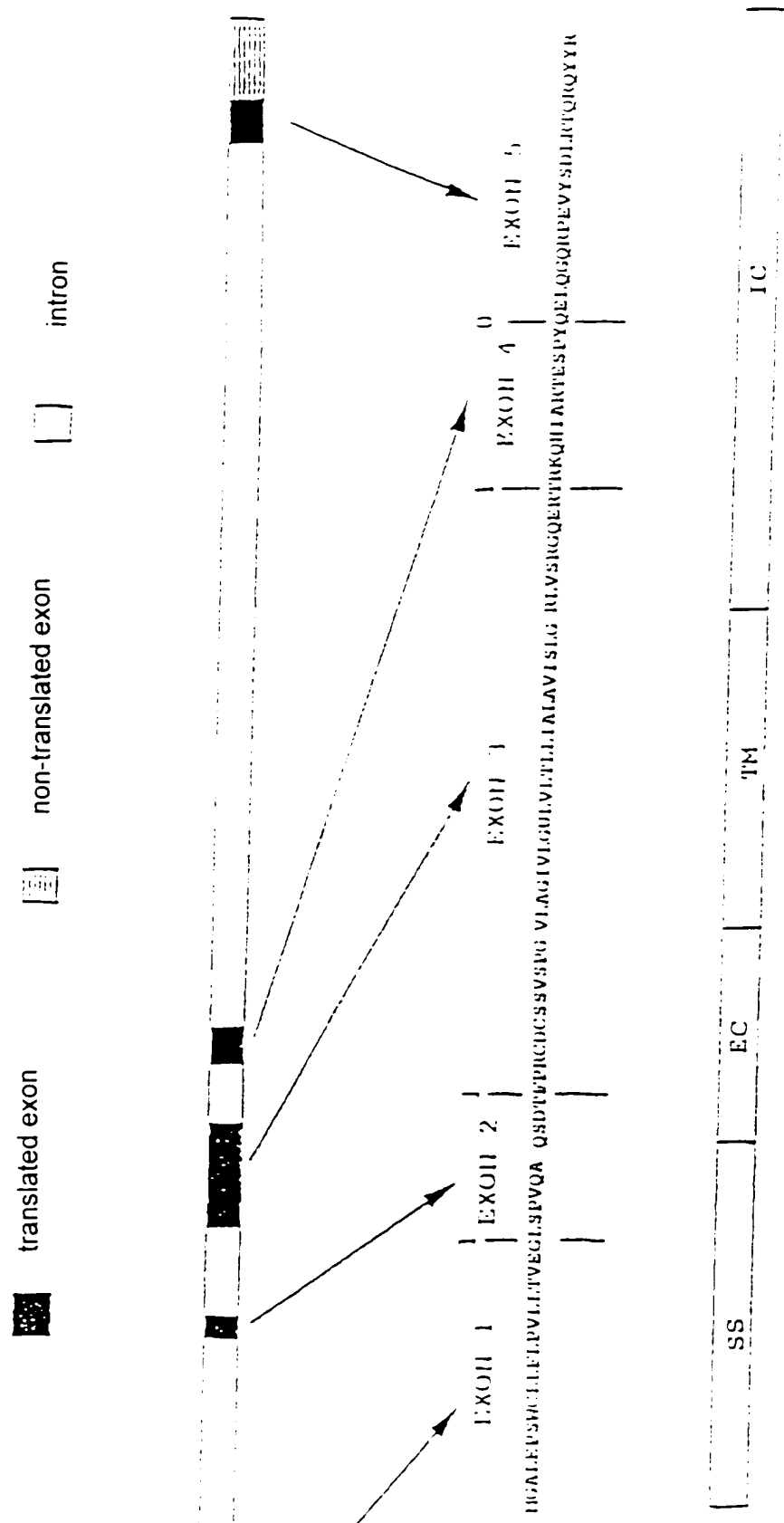


Figure 22

Release of serotonin induced by the p50/KARAP
complex reconstituted in RBL-2H3 cells

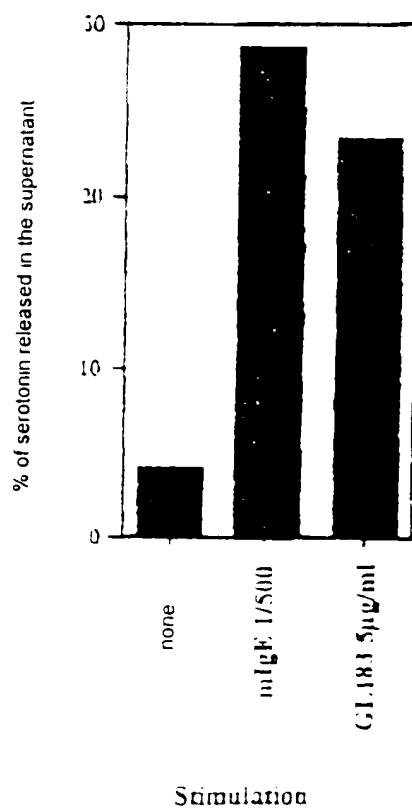


Figure 23

